## SEQUENCE LISTING

<110> Novozymes Biotech, Inc. <120> Microbial Trypsin Mutants Having Chymotrypsin Activity And Nucleic Acids Encoding Same <130> 10211.204-US 60/413,057 <150> 2002-09-24 <151> <160> 23 <170> PatentIn version 3.2 <210> 1 <211> 998 DNA Fusarium oxysporum <213> <400> 1 atcatcaacc actcttcact cttcaactct cctctcttgg atatctatct cttcaccatg 60 gtcaagttcg cttccgtcgt tgcacttgtt gctcccctgg ctgctgccgc tcctcaggag 120 atccccaaca ttgttggtgg cacttctgcc agcgctggcg actttccctt catcgtgagc 180 attagccgca acggtggccc ctggtgtgga ggttctctcc tcaacgccaa caccgtcttg 240 actgctgccc actgcgtttc cggatacgct cagagcggtt tccagattcg tgctggcagt 300 ctgtctcgca cttctggtgg tattacctcc tcgctttcct ccgtcagagt tcaccctagc 360 420 tacageggaa acaacaacga tettgetatt etgaagetet etaetteeat eeeeteegge ggaaacatcg gctatgctcg cctggctgct tccggctctg accctgtcgc tggatcttct 480 540 gccactgttg ctggctgggg cgctacctct gagggcggca gctctactcc cgtcaacctt ctgaaggtta ctgtccctat cgtctctcgt gctacctgcc gagctcagta cggcacctcc 600 gccatcacca accagatgtt ctgtgctggt gtttcttccg gtggcaagga ctcttgccag 660 ggtgacagcg gcggccccat cgtcgacagc tccaacactc ttatcggtgc tgtctcttgg 720 ggtaacggat gtgcccgacc caactactct ggtgtctatg ccagcgttgg tgctctccgc 780 tctttcattg acacctatgc ttaaatacct tgttggaagc gtcgagatgt tccttgaata 840 ttctctagct tgagtcttgg atacgaaacc tgtttgagaa ataggtttca acgagttaag 900 aagatatgag ttgatttcag ttggatctta gtcctggttg ctcgtaatag agcaatctag 960 998

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Ala Gly Asp Phe Pro Phe Ile Val Ser Ile Ser Arg Asn Gly Gly Pro 35 40 45

Trp Cys Gly Gly Ser Leu Leu Asn Ala Asn Thr Val Leu Thr Ala Ala 50 55 60

His Cys Val Ser Gly Tyr Ala Gln Ser Gly Phe Gln Ile Arg Ala Gly 65 70 75 80

Ser Leu Ser Arg Thr Ser Gly Gly Ile Thr Ser Ser Leu Ser Ser Val 85 90 95

Arg Val His Pro Ser Tyr Ser Gly Asn Asn Asn Asp Leu Ala Ile Leu
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Lys Leu Ser Thr Ser Ile Pro Ser Gly Gly Asn Ile Gly Tyr Ala Arg 115 120 125

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Ala Gly Trp Gly Ala Thr Ser Glu Gly Gly Ser Ser Thr Pro Val Asn 145 150 155 160

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Gln Tyr Gly Thr Ser Ala Ile Thr Asn Gln Met Phe Cys Ala Gly Val 180 185 190 Ser Ser Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Ile 195 200 205

Val Asp Ser Ser Asn Thr Leu Ile Gly Ala Val Ser Trp Gly Asn Gly 210 215 220

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Ala Gly Asp Phe Pro Phe Ile Val Ser Ile Ser Arg Asn Gly Gly Pro 35 40 45

Trp Cys Gly Gly Ser Leu Leu Asn Ala Asn Thr Val Leu Thr Ala Ala 50 55 60

His Cys Val Ser Gly Tyr Ala Gln Ser Gly Phe Gln Ile Arg Ala Gly 65 70 75 80

Ser Leu Ser Arg Thr Ser Gly Gly Ile Thr Ser Ser Leu Ser Ser Val 85 90 95

Arg Val His Pro Ser Tyr Ser Gly Asn Asn Asn Asp Leu Ala Ile Leu 100 105 110

Lys Leu Ser Thr Ser Ile Pro Ser Gly Gly Asn Ile Gly Tyr Ala Arg 115 120 125

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